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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/014,743

DATE: 01/02/2002

TIME: 11:28:01

Input Set : A:\Ascii (oct-01).seq.txt
Output Set: N:\CRF3\01022002\J014743.raw

ENTERED

SEQUENCE LISTING

2 (1) GENERAL INFORMATION:

3 (i) APPLICANT: PE Corporation (NY)

4 (ii) TITLE OF INVENTION: ENERGY TRANSFER DYES WITH ENHANCED FLUORESCENCE

5 (iii) NUMBER OF SEQUENCES: 3

6 (iv) CORRESPONDENCE ADDRESS:

7 (A) ADDRESSEE: David J. Weitz, Wilson Sonsini Goodrich
& Rosati

8 (B) STREET: 650 Page Mill Road

9 (C) CITY: Palo Alto

10 (D) STATE: California

11 (E) COUNTRY: USA

12 (F) ZIP: 94304-1050

13 (v) COMPUTER READABLE FORM:

14 (A) MEDIUM TYPE: 3.5 inch diskette

15 (B) COMPUTER: IBM compatible

16 (C) OPERATING SYSTEM: Microsoft Windows 3.1/DOS 5.0

17 (D) SOFTWARE: Word 97

18 ASCII (DOS) TEXT format

19 (vi) CURRENT APPLICATION DATA:

20 (A) APPLICATION NUMBER: US/10/014,743

21 (B) FILING DATE: 29-Oct-2001

22 (C) CLASSIFICATION:

23 (vii) PRIOR APPLICATION DATA:

24 (A) APPLICATION NUMBER: 08/642,330

25 (B) FILING DATE: May 3, 1996

26 (A) APPLICATION NUMBER: 08/672,196

27 (B) FILING DATE: June 27, 1996

28 (A) APPLICATION NUMBER: 08/726,462

29 (B) FILING DATE: October 4, 1996

30 (A) APPLICATION NUMBER: 09/046,203

31 (B) FILING DATE: March 23, 1998

32 (A) APPLICATION NUMBER: 09/272,097

33 (B) FILING DATE: March 18, 1999

34 (viii) ATTORNEY/AGENT INFORMATION:

35 (A) NAME: David J. Weitz

36 (B) REGISTRATION NUMBER: 38,362

37 (C) REFERENCE/DOCKET NUMBER: 16842-776

38 (ix) TELECOMMUNICATION INFORMATION:

39 (A) TELEPHONE: (650) 493-9300

40 (B) TELEFAX: (650) 493-6811

41 (2) INFORMATION FOR SEQ ID NO: 1:

42 (i) SEQUENCE CHARACTERISTICS:

43 (A) LENGTH: 1217 nucleotides

44 (B) TYPE: nucleic acid

45 (C) STRANDEDNESS: single

46 (D) TOPOLOGY: linear

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63 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
 64 GCCAAGCTTG CATGCCTGCA GGTCGACTCT AGAGGATCCC 40
 65 CGGGTACCGA GCTCGAATTG GTAATCATGG TCATAGCTGT 80
 66 TTCCTGTGTG AAATTGTTAT CCGCTCACAA TTCCACACAA 120
 67 CATA CGAGCC GGAAGCATAA AGTGTAAAGC CTGGGGTGCC 160
 68 TAATGAGTGA GCTAACTCAC ATTAATTGCG TTGCGCTCAC 200
 69 TGCCCGCTTT CCAGTCGGGA AACCTGTCGT GCCAGCTGCA 240
 70 TTAATGAATC GGCCAACGCG CGGGGAGAGG CGGTTTGCCT 280
 71 ATTGGGCGCC AGGGTGGTTT TTCTTTTCAC CAGTGAGACG 320
 72 GGCAACAGCT GATTGCCCTT CACCGCCTGG CCCTGAGAGA 360
 73 GTTGCAGCAA GCGGTCCACG CTGGTTTGCC CCAGCAGGCG 400
 74 AAAATCCTGT TTGATGGTGG TTCCGAAATC GGCAAAATCC 440
 75 CTTATAAACTC AAAAGAATAG CCCGAGATAG GGTTGAGTGT 480
 76 TGTCCAGTT TGGAAACAAGA GTCCACTATT AAAGAACGTG 520
 77 GACTCCAACG TCAAAGGGCG AAAAACCGTC TATCAGGGCG 560
 78 ATGGCCCACCT ACGTGAACCA TCACCCAAAT CAAGTTTTT 600
 79 GGGGTCGAGG TGCCGTAAAG CACTAAATCG GAACCTAA 640
 80 GGGAGCCCCC GATTTAGAGC TTGACGGGA AAGCCGGCGA 680
 81 ACGTGGCGAG AAAGGAAGGG AAGAAAGCGA AAGGAGCGGG 720
 82 CGCTAGGGCG CTGGCAAGTG TAGCGGTACAC GCTGCGCGTA 760
 83 ACCACCACAC CCGCCCGCCT TAATGCGCCG CTACAGGGCG 800
 84 CGTACTATGG TTGCTTGAC GAGCACGTAT AACGTGCTTT 840
 85 CCTCGTTGGA ATCAGAGCGG GAGCTAAACA GGAGGCGCAT 880
 86 TAAAGGGATT TTAGACAGGA ACGGTACGCC AGAATCTTGA 920
 87 GAAGTGTGTT TATAATCAGT GAGGCCACCG AGTAAAAGAG 960
 88 TCTGTCCATC ACGCAAATTA ACCGTTGTAG CAATACTTCT 1000
 89 TTGATTAGTA ATAACATCAC TTGCCTGAGT AGAAGAACTC 1040
 90 AACTATCGG CTTGCTGGT AATATCCAGA ACAATATTAC 1080
 91 CGCCAGCCAT TGCAACAGGA AAAACGCTCA TGGAAATACC 1120
 92 TACATTTGA CGCTCAATCG TCTGAAATGG ATTATTTACA 1160
 93 TTGGCAGATT CACCAGTCAC ACGACCAGTA ATAAAAGGGA 1200
 94 CATTCTGGCC AACAGAG 1217

96 (2) INFORMATION FOR SEQ ID NO: 2:

97 (i) SEQUENCE CHARACTERISTICS:
 98 (A) LENGTH: 18 nucleotides
 99 (B) TYPE: nucleic acid
 100 (C) STRANDEDNESS: single
 101 (D) TOPOLOGY: linear

102 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

103 TGTAAAACGA CGGCCAGT 18

106 (2) INFORMATION FOR SEQ ID NO: 3:

107 (i) SEQUENCE CHARACTERISTICS:
 108 (A) LENGTH: 738 nucleotides
 109 (B) TYPE: nucleic acid
 110 (C) STRANDEDNESS: single
 111 (D) TOPOLOGY: linear

112 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

113 ATACGACTCA CTATAGGGCG AATTCGAGCT CGGTACCCGG 40
 114 GGATCCTCTA GAGTCGACCT GCAGGCATGC AAGCTTGAGT 80

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115	ATTCTATAGT	GTCACCTAAA	TAGCTTGGCG	TAATCATGGT	120
116	CATAGCTGTT	TCCTGTGTGA	AATTGTTATC	CGCTCACAAAT	160
117	TCCACACAAC	ATACGAGCCG	GAAGCATAAA	GTGTAAAGCC	200
118	TGGGGTGCCT	AATGAGTGAG	CTAACTCACA	TTAATTGCGT	240
119	TGCGCTCACT	GCCC GCTTTC	CAGTCGGGAA	ACCTGTCGTG	280
120	CCAGCTGCAT	TAATGAATCG	GCCAACGCGC	GGGGAGAGGC	320
121	GGTTTGCGTA	TTGGGCGCTC	TTCCGCTTCC	TCGCTCACTG	360
122	ACTCGCTGCG	CTCGGTCGTT	CGGCTGCGGC	GAGCGGTATC	400
123	AGCTCACTCA	AAGGC GGTTAA	TACGGTTATC	CACAGAACATCA	440
124	GGGGATAACG	CAGGAAAGAA	CATGTGAGCA	AAAGGCCAGC	480
125	AAAAGGCCAG	GAACCGTAAA	AAGGCCCGT	TGCTGGCGTT	520
126	TTTCCATAGG	CTCCGCCCCC	CTGACGAGCA	TCACAAAAAT	560
127	CGACGCTCAA	GTCAGAGGTG	GCGAAACCCG	ACAGGACTAT	600
128	AAAGATACCA	GGCGTTTCCC	CCTGGAAGCT	CCCTCGTGC	640
129	CTCTCCTGTT	CCGACCCCTGC	CGCTTACCGG	ATACCTGTCC	680
130	GCCTTCTCC	CTTCGGGAAG	CGTGGCGCTT	TCTCATAGCT	720
131	CACGCTGTAG	GTATCTCA			738

VERIFICATION SUMMARY
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L:23 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:24 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]